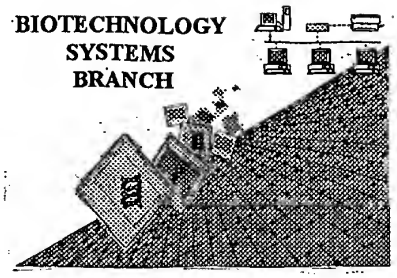
0590

0~~8~~0

2/22

BIOTECHNOLOGY SYSTEMS BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/040,906
Source: OIPE
Date Processed by STIC: 1/23/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY** or,
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 3.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom, including:

1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm**>, **EFS Submission User Manual** - ePAVE)
2. U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. **Hand Carry directly to:**
   U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
   Or
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. **Federal Express, United Parcel Service, or other delivery service to:** U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 10/040,906

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1	Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2	Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3	Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4	Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5	Variable Length	Sequence(s)____contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6	PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7	Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8	Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9	Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10	Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
11	Use of <220>	Sequence(s) 8 missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12	PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13	Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/040,906

DATE: 01/23/2002
TIME: 19:05:45

Input Set : A:\wnew2asus2.st25.txt
Output Set: N:\CRF3\01232002\J040906.raw

p6

Does Not Comply
Corrected Diskette Needed

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 2 <110> APPLICANT: Arnaut, Greta
 3       Boets, Annemie
 4       Vanneste, Stijn
 5       Van Rie, Jeroen
 6       Van Houdt, Sara
 8 <120> TITLE OF INVENTION: Novel Bacillus thuringiensis insecticidal proteins
10 <130> FILE REFERENCE: 58764.000036
12 <140> CURRENT APPLICATION NUMBER: US/10/040,906
C-> 12 <141> CURRENT FILING DATE: 2002-01-09
12 <150> PRIOR APPLICATION NUMBER: US 09/756,296
13 <151> PRIOR FILING DATE: 2001-01-09
15 <160> NUMBER OF SEQ ID NOS: 9
17 <170> SOFTWARE: PatentIn version 3.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 1899
21 <212> TYPE: DNA
22 <213> ORGANISM: Bacillus thuringiensis
24 <220> FEATURE:
25 <221> NAME/KEY: CDS
26 <222> LOCATION: (1)..(1896)
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32 1               5                  10                  15
34 aat gta gtg gcc cat gat cca ttt agt ttt gag cat aaa tca tta gat      96
35 Asn Val Val Ala His Asp Pro Phe Ser Phe Glu His Lys Ser Leu Asp
36             20                  25                  30
38 acc atc cga aaa gaa tgg atg gag tgg aaa aga aca gat cat agt tta     144
39 Thr Ile Arg Lys Glu Trp Met Glu Trp Lys Arg Thr Asp His Ser Leu
40         35                  40                  45
42 tat gta gct cct ata gtc gga act gtt tct agc ttt ctg cta aag aag     192
43 Tyr Val Ala Pro Ile Val Gly Thr Val Ser Ser Phe Leu Leu Lys Lys
44     50                  55                  60
46 gtg ggg agt ctt att gga aaa agg ata ttg agt gaa tta tgg ggg tta     240
47 Val Gly Ser Leu Ile Gly Lys Arg Ile Leu Ser Glu Leu Trp Gly Leu
48 65                  70                  75                  80
50 ata ttt cct agt ggt agc aca aat cta atg caa gat att tta agg gag     288
51 Ile Phe Pro Ser Gly Ser Thr Asn Leu Met Gln Asp Ile Leu Arg Glu
52                 85                  90                  95
54 aca gaa caa ttc cta aat caa aga ctt aat aca gac act ctt gcc cgt     336
55 Thr Glu Gln Phe Leu Asn Gln Arg Leu Asn Thr Asp Thr Leu Ala Arg
56             100                 105                 110
58 gta aat gcg gaa ttg gaa ggg ctg caa gcg aat ata agg gag ttt aat     384
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/040,906

DATE: 01/23/2002
TIME: 19:05:45

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Output Set: N:\CRF3\01232002\J040906.raw

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60         115                 120                 125
62 caa caa gta gat aat ttt tta aat cct act caa aac cct gtt cct tta      432
63 Gln Gln Val Asp Asn Phe Leu Asn Pro Thr Gln Asn Pro Val Pro Leu
64     130                 135                 140
66 tca ata act tct tca gtt aat aca atg cag caa tta ttt cta aat aga      480
67 Ser Ile Thr Ser Ser Val Asn Thr Met Gln Gln Leu Phe Leu Asn Arg
68 145                 150                 155                 160
70 tta ccc cag ttc cgt gtg caa gga tac caa ctg tta tta tta cct tta      528
71 Leu Pro Gln Phe Arg Val Gln Gly Tyr Gln Leu Leu Leu Leu Pro Leu
72                 165                 170                 175
74 ttt gca cag gca gcc aat atg cat ctt tct ttt att aga gat gtt gtt      576
75 Phe Ala Gln Ala Ala Asn Met His Leu Ser Phe Ile Arg Asp Val Val
76             180                 185                 190
78 ctc aat gca gat gaa tgg gga att tca gca gca aca tta cgt acg tat      624
79 Leu Asn Ala Asp Glu Trp Gly Ile Ser Ala Ala Thr Leu Arg Thr Tyr
80         195                 200                 205
82 caa aat tat ctg aaa aat tat aca aca gag tac tct aat tat tgt ata      672
83 Gln Asn Tyr Leu Lys Asn Tyr Thr Thr Glu Tyr Ser Asn Tyr Cys Ile
84     210                 215                 220
86 aat acg tat caa act gcg ttt aga ggt tta aac acc cgt tta cac gat      720
87 Asn Thr Tyr Gln Thr Ala Phe Arg Gly Leu Asn Thr Arg Leu His Asp
88 225                 230                 235                 240
90 atg tta gaa ttt aga aca tat atg ttt tta aat gta ttt gaa tat gta      768
91 Met Leu Glu Phe Arg Thr Tyr Met Phe Leu Asn Val Phe Glu Tyr Val
92                 245                 250                 255
94 tct atc tgg tcg ttg ttt aaa tat caa agc ctt cta gta tct tct ggc      816
95 Ser Ile Trp Ser Leu Phe Lys Tyr Gln Ser Leu Leu Val Ser Ser Gly
96             260                 265                 270
98 gct aat tta tat gca agc ggt agt gga cca cag cag act caa tca ttt      864
99 Ala Asn Leu Tyr Ala Ser Gly Ser Gly Pro Gln Gln Thr Gln Ser Phe
100         275                 280                 285
102 act tca caa gac tgg cca ttt tta tat tct ctt ttc caa gtt aat tca      912
103 Thr Ser Gln Asp Trp Pro Phe Leu Tyr Ser Leu Phe Gln Val Asn Ser
104     290                 295                 300
106 aat tat gtg tta aat ggc ttt agt ggc gct aga ctt acg cag act ttc      960
107 Asn Tyr Val Leu Asn Gly Phe Ser Gly Ala Arg Leu Thr Gln Thr Phe
108 305                 310                 315                 320
110 cct aat att ggt ggt tta cct ggt act act aca act cac gca ttg ctt     1008
111 Pro Asn Ile Gly Gly Leu Pro Gly Thr Thr Thr Thr His Ala Leu Leu
112                 325                 330                 335
114 gcg gca agg gtc aat tac agt gga gga gtt tcg tct ggt gat ata ggc     1056
115 Ala Ala Arg Val Asn Tyr Ser Gly Gly Val Ser Ser Gly Asp Ile Gly
116             340                 345                 350
118 gct gtg ttt aat caa aat ttt agt tgt agc aca ttt ctc cca cct ttg     1104
119 Ala Val Phe Asn Gln Asn Phe Ser Cys Ser Thr Phe Leu Pro Pro Leu
120         355                 360                 365
122 tta aca cca ttt gtt agg agt tgg cta gat tca ggt tca gat cga ggg     1152
123 Leu Thr Pro Phe Val Arg Ser Trp Leu Asp Ser Gly Ser Asp Arg Gly
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/040,906

DATE: 01/23/2002
TIME: 19:05:45

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Output Set: N:\CRF3\01232002\J040906.raw

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124     370                 375                 380
126 ggt gtt aat acc gtt aca aat tgg caa aca gaa tcg ttt gag tca act     1200
127 Gly Val Asn Thr Val Thr Asn Trp Gln Thr Glu Ser Phe Glu Ser Thr
128 385                 390                 395                 400
130 tta ggt tta agg tgt ggt gct ttt aca gct cgt ggt aat tca aac tat     1248
131 Leu Gly Leu Arg Cys Gly Ala Phe Thr Ala Arg Gly Asn Ser Asn Tyr
132                 405                 410                 415
134 ttc cca gat tat ttt atc cgt aat att tca gga gtt cct tta gtt gtt     1296
135 Phe Pro Asp Tyr Phe Ile Arg Asn Ile Ser Gly Val Pro Leu Val Val
136             420                 425                 430
138 aga aat gaa gat tta aga aga ccg tta cac tat aat gaa ata aga aat     1344
139 Arg Asn Glu Asp Leu Arg Arg Pro Leu His Tyr Asn Glu Ile Arg Asn
140         435                 440                 445
142 ata gaa agt cct tca gga aca cct ggt gga tta cga gct tat atg gta     1392
143 Ile Glu Ser Pro Ser Gly Thr Pro Gly Gly Leu Arg Ala Tyr Met Val
144     450                 455                 460
146 tct gtg cat aat aga aaa aat aat atc tat gcc gtg cat gaa aat ggt     1440
147 Ser Val His Asn Arg Lys Asn Asn Ile Tyr Ala Val His Glu Asn Gly
148 465                 470                 475                 480
150 act atg att cat tta gcg ccg gaa gat tat aca gga ttc acc ata tcg     1488
151 Thr Met Ile His Leu Ala Pro Glu Asp Tyr Thr Gly Phe Thr Ile Ser
152                 485                 490                 495
154 ccg ata cat gca act caa gtg aat aat caa acg cga aca ttt att tct     1536
155 Pro Ile His Ala Thr Gln Val Asn Asn Gln Thr Arg Thr Phe Ile Ser
156             500                 505                 510
158 gaa aaa ttt gga aat caa ggt gat tcc tta aga ttt gaa caa agc aac     1584
159 Glu Lys Phe Gly Asn Gln Gly Asp Ser Leu Arg Phe Glu Gln Ser Asn
160         515                 520                 525
162 acg aca gca cgt tat aca ctt aga gga aat gga aat agt tac aat ctt     1632
163 Thr Thr Ala Arg Tyr Thr Leu Arg Gly Asn Gly Asn Ser Tyr Asn Leu
164     530                 535                 540
166 tat tta aga gta tct tca cta gga aat tcc act att cga gtt act ata     1680
167 Tyr Leu Arg Val Ser Ser Leu Gly Asn Ser Thr Ile Arg Val Thr Ile
168 545                 550                 555                 560
170 aac ggt agg gtt tat act gct tca aat gtt aat act act aca aat aac     1728
171 Asn Gly Arg Val Tyr Thr Ala Ser Asn Val Asn Thr Thr Thr Asn Asn
172                 565                 570                 575
174 gat gga gtt aat gat aat ggc gct cgt ttt tta gat att aat atg ggt     1776
175 Asp Gly Val Asn Asp Asn Gly Ala Arg Phe Leu Asp Ile Asn Met Gly
176             580                 585                 590
178 aat gta gta gca agt gat aat act aat gta ccg tta gat ata aat gtg     1824
179 Asn Val Val Ala Ser Asp Asn Thr Asn Val Pro Leu Asp Ile Asn Val
180         595                 600                 605
182 aca ttt aac tcc ggt act caa ttt gag ctt atg aat att atg ttt gtt     1872
183 Thr Phe Asn Ser Gly Thr Gln Phe Glu Leu Met Asn Ile Met Phe Val
184     610                 615                 620
186 cca act aat ctt cca cca ata tat taa                               1899
187 Pro Thr Asn Leu Pro Pro Ile Tyr
188 625                 630
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RAW SEQUENCE LISTING DATE: 01/23/2002
PATENT APPLICATION: US/10/040,906 TIME: 19:05:45

Input Set : A:\wnew2asus2.st25.txt
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192 <211> LENGTH: 632
193 <212> TYPE: PRT
194 <213> ORGANISM: Bacillus thuringiensis
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199 1               5                  10                  15
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203             20                  25                  30
206 Thr Ile Arg Lys Glu Trp Met Glu Trp Lys Arg Thr Asp His Ser Leu
207         35                  40                  45
210 Tyr Val Ala Pro Ile Val Gly Thr Val Ser Ser Phe Leu Leu Lys Lys
211     50                  55                  60
214 Val Gly Ser Leu Ile Gly Lys Arg Ile Leu Ser Glu Leu Trp Gly Leu
215 65                  70                  75                  80
218 Ile Phe Pro Ser Gly Ser Thr Asn Leu Met Gln Asp Ile Leu Arg Glu
219                 85                  90                  95
222 Thr Glu Gln Phe Leu Asn Gln Arg Leu Asn Thr Asp Thr Leu Ala Arg
223             100                 105                 110
226 Val Asn Ala Glu Leu Glu Gly Leu Gln Ala Asn Ile Arg Glu Phe Asn
227         115                 120                 125
230 Gln Gln Val Asp Asn Phe Leu Asn Pro Thr Gln Asn Pro Val Pro Leu
231     130                 135                 140
234 Ser Ile Thr Ser Ser Val Asn Thr Met Gln Gln Leu Phe Leu Asn Arg
235 145                 150                 155                 160
238 Leu Pro Gln Phe Arg Val Gln Gly Tyr Gln Leu Leu Leu Leu Pro Leu
239                 165                 170                 175
242 Phe Ala Gln Ala Ala Asn Met His Leu Ser Phe Ile Arg Asp Val Val
243             180                 185                 190
246 Leu Asn Ala Asp Glu Trp Gly Ile Ser Ala Ala Thr Leu Arg Thr Tyr
247         195                 200                 205
250 Gln Asn Tyr Leu Lys Asn Tyr Thr Thr Glu Tyr Ser Asn Tyr Cys Ile
251     210                 215                 220
254 Asn Thr Tyr Gln Thr Ala Phe Arg Gly Leu Asn Thr Arg Leu His Asp
255 225                 230                 235                 240
258 Met Leu Glu Phe Arg Thr Tyr Met Phe Leu Asn Val Phe Glu Tyr Val
259                 245                 250                 255
262 Ser Ile Trp Ser Leu Phe Lys Tyr Gln Ser Leu Leu Val Ser Ser Gly
263             260                 265                 270
266 Ala Asn Leu Tyr Ala Ser Gly Ser Gly Pro Gln Gln Thr Gln Ser Phe
267         275                 280                 285
270 Thr Ser Gln Asp Trp Pro Phe Leu Tyr Ser Leu Phe Gln Val Asn Ser
271     290                 295                 300
274 Asn Tyr Val Leu Asn Gly Phe Ser Gly Ala Arg Leu Thr Gln Thr Phe
275 305                 310                 315                 320
278 Pro Asn Ile Gly Gly Leu Pro Gly Thr Thr Thr Thr His Ala Leu Leu
279                 325                 330                 335
282 Ala Ala Arg Val Asn Tyr Ser Gly Gly Val Ser Ser Gly Asp Ile Gly
283             340                 345                 350
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RAW SEQUENCE LISTING DATE: 01/23/2002
PATENT APPLICATION: US/10/040,906 TIME: 19:05:45

Input Set : A:\wnew2asus2.st25.txt
Output Set: N:\CRF3\01232002\J040906.raw

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286 Ala Val Phe Asn Gln Asn Phe Ser Cys Ser Thr Phe Leu Pro Pro Leu
287         355                 360                 365
290 Leu Thr Pro Phe Val Arg Ser Trp Leu Asp Ser Gly Ser Asp Arg Gly
291     370                 375                 380
294 Gly Val Asn Thr Val Thr Asn Trp Gln Thr Glu Ser Phe Glu Ser Thr
295 385                 390                 395                 400
298 Leu Gly Leu Arg Cys Gly Ala Phe Thr Ala Arg Gly Asn Ser Asn Tyr
299                 405                 410                 415
302 Phe Pro Asp Tyr Phe Ile Arg Asn Ile Ser Gly Val Pro Leu Val Val
303             420                 425                 430
306 Arg Asn Glu Asp Leu Arg Arg Pro Leu His Tyr Asn Glu Ile Arg Asn
307         435                 440                 445
310 Ile Glu Ser Pro Ser Gly Thr Pro Gly Gly Leu Arg Ala Tyr Met Val
311     450                 455                 460
314 Ser Val His Asn Arg Lys Asn Asn Ile Tyr Ala Val His Glu Asn Gly
315 465                 470                 475                 480
318 Thr Met Ile His Leu Ala Pro Glu Asp Tyr Thr Gly Phe Thr Ile Ser
319                 485                 490                 495
322 Pro Ile His Ala Thr Gln Val Asn Asn Gln Thr Arg Thr Phe Ile Ser
323             500                 505                 510
326 Glu Lys Phe Gly Asn Gln Gly Asp Ser Leu Arg Phe Glu Gln Ser Asn
327         515                 520                 525
330 Thr Thr Ala Arg Tyr Thr Leu Arg Gly Asn Gly Asn Ser Tyr Asn Leu
331     530                 535                 540
334 Tyr Leu Arg Val Ser Ser Leu Gly Asn Ser Thr Ile Arg Val Thr Ile
335 545                 550                 555                 560
338 Asn Gly Arg Val Tyr Thr Ala Ser Asn Val Asn Thr Thr Thr Asn Asn
339                 565                 570                 575
342 Asp Gly Val Asn Asp Asn Gly Ala Arg Phe Leu Asp Ile Asn Met Gly
343             580                 585                 590
346 Asn Val Val Ala Ser Asp Asn Thr Asn Val Pro Leu Asp Ile Asn Val
347         595                 600                 605
350 Thr Phe Asn Ser Gly Thr Gln Phe Glu Leu Met Asn Ile Met Phe Val
351     610                 615                 620
354 Pro Thr Asn Leu Pro Pro Ile Tyr
355 625                 630
358 <210> SEQ ID NO: 3
359 <211> LENGTH: 1899
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361 <213> ORGANISM: Bacillus thuringiensis
363 <220> FEATURE:
364 <221> NAME/KEY: CDS
365 <222> LOCATION: (1)..(1896)
367 <400> SEQUENCE: 3
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371 1               5                   10                  15
373 aat gta gtg gct cat gat cca ttt agt ttt caa cat aaa tca tta gat       96
374 Asn Val Val Ala His Asp Pro Phe Ser Phe Gln His Lys Ser Leu Asp
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<210> 8
<211> 633
<212> PRT
<213> Artificial Sequence

see item 11 on Error Summary Sheet

<400> 8

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/040,906

DATE: 01/23/2002
TIME: 19:05:46

Input Set : A:\wnew2asus2.st25.txt
Output Set: N:\CRF3\01232002\J040906.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1217 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1217 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: